

SEQUENCE LISTING

<110> KAKKIS, EMIL D.

<120> METHODS FOR TREATING DISEASES CAUSED BY DEFICIENCIES OF RECOMBINANT ALPHA-L-IDURONIDINASE

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<151> 2000-11-09

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Met

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Leu Ala Pro Pro Val Ala Pro Ala Glu Ala Pro His Leu Val His	20	25	30	
ctg gcc gcg ccc ccc gtg gcc ccg gag gcc ccg cac ctg gtg cat Leu Ala Ala Pro Pro Val Ala Pro Ala Glu Ala Pro His Leu Val His	35	40	45	1656
gtg gac gcg gcc cgc gcg ctg tgg ccc ctg ccg ccg ttc tgg agg agc Val Asp Ala Ala Arg Ala Leu Trp Pro Leu Arg Arg Phe Trp Arg Ser	35	40	45	1704
aca ggc ttc tgc ccc ccg ctg cca cac agc cag gct gac cag tac gtg Thr Gly Phe Cys Pro Pro Leu Pro His Ser Gln Ala Asp Gln Tyr Val	50	55	60	1752
a	70	75	80	
ctc agc tgg gac cag cag ctc aac ctc gcc tat gtg ggc gcc gtc cct Leu Ser Trp Asp Gln Gln Leu Asn Leu Ala Tyr Val Gly Ala Val Pro				1800
cac cgc ggc atc aag cag gtc cgg acc cac tgg ctg ctg gag ctt gtc His Arg Gly Ile Lys Gln Val Arg Thr His Trp Leu Leu Glu Leu Val	85	90	95	1848
acc acc agg ggg tcc act gga cgg ggc ctg agc tac aac ttc acc cac Thr Thr Arg Gly Ser Thr Gly Arg Gly Leu Ser Tyr Asn Phe Thr His	100	105	110	1896
ctg gac ggg tac ctg gac ctt ctc agg gag aac cag ctc ggg ttt gag Leu Asp Gly Tyr Leu Asp Leu Leu Arg Glu Asn Gln Leu Gly Phe Glu	115	120	125	1944
ctg atg ggc agc gcc tcg ggc cac ttc act gac ttt gag gac aag cag Leu Met Gly Ser Ala Ser Gly His Phe Thr Asp Phe Glu Asp Lys Gln	130	135	140	1992
cag gtg ttt gag tgg aag gac ttg gtc tcc agc ctg gcc agg aga tac Gln Val Phe Glu Trp Lys Asp Leu Val Ser Ser Leu Ala Arg Arg Tyr	150	155	160	2040
atc ggt agg tac gga ctg gcg cat gtt tcc aag tgg aac ttc gag acg Ile Gly Arg Tyr Gly Leu Ala His Val Ser Lys Trp Asn Phe Glu Thr	165	170	175	2088
tgg aat gag cca gac cac cac gac ttt gac aac gtc tcc atg acc atg Trp Asn Glu Pro Asp His His Asp Phe Asp Asn Val Ser Met Thr Met	180	185	190	2136
caa ggc ttc ctg aac tac tac gat gcc tgc tcg gag ggt ctg cgc gcc Gln Gly Phe Leu Asn Tyr Tyr Asp Ala Cys Ser Glu Gly Leu Arg Ala	195	200	205	2184
gcc agc ccc gcc ctg cgg ctg gga ggc ccc ggc gac tcc ttc cac acc Ala Ser Pro Ala Leu Arg Leu Gly Gly Pro Gly Asp Ser Phe His Thr	210	215	220	2232
c	225			
cca ccg cga tcc ccc ctg agc tgg ggc ctc ctg cgc cac tgc cac gac Pro Pro Arg Ser Pro Leu Ser Trp Gly Leu Leu Arg His Cys His Asp	230	235	240	2280

ggt acc aac ttc ttc act ggg gag ggc gtc cgg ctg gac tac atc Gly Thr Asn Phe Phe Thr Gly Glu Ala Gly Val Arg Leu Asp Tyr Ile	245	250	255	2328
tcc ctc cac agg aag ggt gcg cgc agc tcc atc tcc atc ctg gag cag Ser Leu His Arg Lys Gly Ala Arg Ser Ser Ile Ser Ile Leu Glu Gln	260	265	270	2376
gag aag gtc gtc gcg cag cag cag ctc ttc ccc aag ttc gcg Glu Lys Val Val Ala Gln Ile Arg Gln Leu Phe Pro Lys Phe Ala	275	280	285	2424
gac acc ccc att tac aac gac gag gcg gac ccc ctg gtc ggc tgg tcc Asp Thr Pro Ile Tyr Asn Asp Glu Ala Asp Pro Leu Val Gly Trp Ser	290	295	300	2472
ctg cca cag ccg tgg agg gcg gac gtc acc tac gcg gcc atg gtc Leu Pro Gln Pro Trp Arg Ala Asp Val Thr Tyr Ala Ala Met Val Val	310	315	320	2520
aag gtc atc gcg cag cat cag aac ctg cta ctg gcc aac acc acc tcc Lys Val Ile Ala Gln His Gln Asn Leu Leu Ala Asn Thr Thr Ser	325	330	335	2568
gcc ttc ccc tac gcg ctc ctg agc aac gac aat gcc ttc ctg agc tac Ala Phe Pro Tyr Ala Leu Leu Ser Asn Asp Asn Ala Phe Leu Ser Tyr	340	345	350	2616
cac ccg cac ccc ttc gcg cag cgc acg ctc acc ggc cgc ttc cag gtc His Pro His Pro Phe Ala Gln Arg Thr Leu Thr Ala Arg Phe Gln Val	355	360	365	2664
aac aac acc cgc ccg ccg cac gtc cag ctg ttg cgc aag ccg gtc Asn Asn Thr Arg Pro Pro His Val Gln Leu Leu Arg Lys Pro Val Leu	370	375	380	2712
acg gcc atg ggg ctg ctg gcg ctg ctg gat gag gag cag ctc tgg gcc Thr Ala Met Gly Leu Leu Ala Leu Leu Asp Glu Glu Gln Leu Trp Ala	390	395	400	2760
gaa gtg tcg cag gcc ggg acc gtc ctg gac agc aac cac acg gtc ggc Glu Val Ser Gln Ala Gly Thr Val Leu Asp Ser Asn His Thr Val Gly	405	410	415	2808
gtc ctg gcc agc gcc cac cgc ccc cag ggc ccg gcc gac gcc tgg cgc Val Leu Ala Ser Ala His Arg Pro Gln Gly Pro Ala Asp Ala Trp Arg	420	425	430	2856
gcc gcg gtg ctg atc tac gcg agc gac gac acc cgc gcc cac ccc aac Ala Ala Val Leu Ile Tyr Ala Ser Asp Asp Thr Arg Ala His Pro Asn	435	440	445	2904
cgc agc gtc gcg gtc acc ctg cgg ctg cgc ggg gtc ccc ccc ggc ccg Arg Ser Val Ala Val Thr Leu Arg Leu Arg Gly Val Pro Pro Gly Pro	450	455	460	2952
ggc ctg gtc tac gtc acg cgc tac ctg gac aac ggg ctc tgc agc ccc Gly Leu Val Tyr Val Thr Arg Tyr Leu Asp Asn Gly Leu Cys Ser Pro	470	475	480	3000
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Asp Gly Glu Trp Arg Arg Leu Gly Arg Pro Val Phe Pro Thr Ala Glu		
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Gln Phe Arg Arg * Arg Ala Ala Glu Asp Pro Val Ala Ala Ala Pro		
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cgc ccc tta ccc gcc ggc ggc cgc ctg agg ctg cgc ccc gcg ctg cgg		3144
Arg Pro Leu Pro Ala Gly Gly Arg Leu Arg Leu Arg Pro Ala Leu Arg		
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ctg cgg tcg ctt ttg ctg gtg cac gtg tgt gcg cgc ccc gag aag ccg		3192
Leu Pro Ser Leu Leu Leu Val His Val Cys Ala Arg Pro Glu Lys Pro		
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ccc ggg cag gtc acg cgg ctc cgc gcc ctg ccc ctg acc caa ggg cag		3240
Pro Gly Gln Val Thr Arg Leu Arg Ala Leu Pro Leu Thr Gln Gly Gln		
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560		
ctg gtt ctg gtc tgg tcg gat gaa cac gtg ggc tcc aag tgc ctg tgg		3288
Leu Val Leu Val Trp Ser Asp Glu His Val Gly Ser Lys Cys Leu Trp		
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aca tac gag atc cag ttc tct cag gac ggt aag ggc tac acc ccc gtc		3336
Thr Tyr Glu Ile Gln Phe Ser Gln Asp Gly Lys Ala Tyr Thr Pro Val		
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Ser Arg Lys Pro Ser Thr Phe Asn Leu Phe Val Phe Ser Pro Asp Thr		
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Gly Ala Val Ser Gly Ser Tyr Arg Val Arg Ala Leu Asp Tyr Trp Ala		
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Pro Arg Gly Pro Pro Ser Pro Gly Asn Pro		
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His	Val	Asp	Ala	Ala	Arg	Ala	Leu	Trp	Pro	Leu	Arg	Arg	Phe	Trp	Arg
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Pro	His	Arg	Gly	Ile	Lys	Gln	Val	Arg	Thr	His	Trp	Leu	Leu	Glu	Leu
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Val	Thr	Thr	Arg	Gly	Ser	Thr	Gly	Arg	Gly	Leu	Ser	Tyr	Asn	Phe	Thr
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His	Leu	Asp	Gly	Tyr	Leu	Asp	Leu	Leu	Arg	Glu	Asn	Gln	Leu	Gly	Phe
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Glu	Leu	Met	Gly	Ser	Ala	Ser	Gly	His	Phe	Thr	Asp	Phe	Glu	Asp	Lys
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Tyr	Ile	Gly	Arg	Tyr	Gly	Leu	Ala	His	Val	Ser	Lys	Trp	Asn	Phe	Glu
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Thr	Trp	Asn	Glu	Pro	Asp	His	His	Asp	Phe	Asp	Asn	Val	Ser	Met	Thr
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Ala	Ala	Ser	Pro	Ala	Leu	Arg	Leu	Gly	Gly	Pro	Gly	Asp	Ser	Phe	His

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